RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/736,892
Source: IFW0
Date Processed by STIC: 3/15/05

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RAW SEQUENCE LISTING DATE: 03/15/2005
PATENT APPLICATION: **US/10/736,892** TIME: 14:23:18

Input Set : **D:\US Utility 50229-424 Sequence Listing.txt**
Output Set: **N:\CRF4\03152005\J736892.raw**

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      3 <110> APPLICANT: University of Kentucky Research Foundation
      4       JI, Tai
      5       JI, Inhae
      7 <120> TITLE OF INVENTION: GENES AND AGENTS TO REGULATE FOLLICULAR DEVELOPMENT,
OVULATION
      8       CYCLE AND STERIODOGENESIS
     10 <130> FILE REFERENCE: 050229-0424
     12 <140> CURRENT APPLICATION NUMBER: 10/736,892
     13 <141> CURRENT FILING DATE: 2003-12-17
     15 <150> PRIOR APPLICATION NUMBER: 60/437,729
     16 <151> PRIOR FILING DATE: 2003-01-03
     18 <160> NUMBER OF SEQ ID NOS: 13
     20 <170> SOFTWARE: PatentIn version 3.3
     22 <210> SEQ ID NO: 1
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     24 <212> TYPE: DNA
     25 <213> ORGANISM: Artificial Sequence
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     28 <223> OTHER INFORMATION: Chemically synthesized
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     39 <220> FEATURE:
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     49 <213> ORGANISM: Artificial Sequence
     51 <220> FEATURE:
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     55 gctttccctc tgttgaccca c                                              21
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     61 <213> ORGANISM: Artificial Sequence
     63 <220> FEATURE:
     64 <223> OTHER INFORMATION: Chemically synthesized
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72 <212> TYPE: DNA
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75 <220> FEATURE:
76 <223> OTHER INFORMATION: Chemically synthesized
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83 <211> LENGTH: 21
84 <212> TYPE: DNA
85 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: Chemically synthesized
90 <400> SEQUENCE: 6
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95 <211> LENGTH: 1099
96 <212> TYPE: DNA
97 <213> ORGANISM: Rattus norvegicus
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102 tccgcagctc cagcctcctc atcccgcccc tggaggccgc actggccaac ttctccaaag 120
104 gtcccgacgg aggggtcatg caaccggtgc gcaccgtggt gcctgtggcc aagcaccgag 180
106 gcttcttggg agtcatgcca gcctacagtg ccgctgagga tgcactcacc accaagttag 240
108 tcaccttcta tgagggccac agcaacaatg ctgtcccctc ccaccaggca tcagtgcttc 300
110 tctttgatcc cagcaatggt tccctgctgg cggtcatgga tggaaatgtc ataactgcaa 360
112 agaggacagc agccgtctct gccatcgcca ccaagttttt gaagccccca ggcagtgatg 420
114 tgctgtgcat tcttggggct ggggtccagg cgtacagtca ctatgagatc ttcacagaac 480
116 agttctcctt caaggaggtg agaatgtgga accgcaccag ggaaaatgct gagaagtttg 540
118 caagctcagt gcagggagat gttcgggtct gttcatcagt gcaggaggct gtgacaggtg 600
120 ccgatgtcat catcacagtc accatggcaa cggagcccat tttatttggt gaatgggtga 660
122 agcccggggc tcacatcaat gctgttggag ccagtagacc tgactggcga gaactggatg 720
124 acgagctcat gaagcaagca gtgctgtatg tggactcccg ggaggctgcc ctaaaggagt 780
126 caggagatgt tctgttgtca ggggctgaca tctttgctga gcttggagaa gtggtttcag 840
128 gagcgaagcc tgcatactgt gagaagacca cggtgttcaa gtctttgggg atggcagtgg 900
130 aggacctggt cgcagccaaa ttagtgtacg attcgtggtc atctggcaag tgagcagaag 960
132 gagctgtgcc tgggctggat ggacgtcacg gctcaaacgc tggctcagtg tctagatcaa 1020
134 aggaggccta gtccccagtg aacgggagtg agagtcactc ataagtattg acatccctat 1080
136 tcatgtttgt ggttggata 1099
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140 <211> LENGTH: 313
141 <212> TYPE: PRT
142 <213> ORGANISM: Rattus norvegicus
144 <400> SEQUENCE: 8
146 Met Arg Arg Ala Pro Ala Phe Leu Ser Ala Asp Glu Val Gln Asp His
147 1 5 10 15
150 Leu Arg Ser Ser Ser Leu Leu Ile Pro Pro Leu Glu Ala Ala Leu Ala

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151              20                  25                  30
154 Asn Phe Ser Lys Gly Pro Asp Gly Gly Val Met Gln Pro Val Arg Thr
155          35                  40                  45
158 Val Val Pro Val Ala Lys His Arg Gly Phe Leu Gly Val Met Pro Ala
159      50                  55                  60
162 Tyr Ser Ala Ala Glu Asp Ala Leu Thr Thr Lys Leu Val Thr Phe Tyr
163 65                  70                  75                  80
166 Glu Gly His Ser Asn Asn Ala Val Pro Ser His Gln Ala Ser Val Leu
167                 85                  90                  95
170 Leu Phe Asp Pro Ser Asn Gly Ser Leu Leu Ala Val Met Asp Gly Asn
171             100                 105                 110
174 Val Ile Thr Ala Lys Arg Thr Ala Ala Val Ser Ala Ile Ala Thr Lys
175          115                 120                 125
178 Phe Leu Lys Pro Pro Gly Ser Asp Val Leu Cys Ile Leu Gly Ala Gly
179      130                 135                 140
182 Val Gln Ala Tyr Ser His Tyr Glu Ile Phe Thr Glu Gln Phe Ser Phe
183 145                 150                 155                 160
186 Lys Glu Val Arg Met Trp Asn Arg Thr Arg Glu Asn Ala Glu Lys Phe
187                 165                 170                 175
190 Ala Ser Ser Val Gln Gly Asp Val Arg Val Cys Ser Ser Val Gln Glu
191             180                 185                 190
194 Ala Val Thr Gly Ala Asp Val Ile Ile Thr Val Thr Met Ala Thr Glu
195          195                 200                 205
198 Pro Ile Leu Phe Gly Glu Trp Val Lys Pro Gly Ala His Ile Asn Ala
199      210                 215                 220
202 Val Gly Ala Ser Arg Pro Asp Trp Arg Glu Leu Asp Asp Glu Leu Met
203 225                 230                 235                 240
206 Lys Gln Ala Val Leu Tyr Val Asp Ser Arg Glu Ala Ala Leu Lys Glu
207                 245                 250                 255
210 Ser Gly Asp Val Leu Leu Ser Gly Ala Asp Ile Phe Ala Glu Leu Gly
211             260                 265                 270
214 Glu Val Val Ser Gly Ala Lys Pro Ala Tyr Cys Glu Lys Thr Thr Val
215          275                 280                 285
218 Phe Lys Ser Leu Gly Met Ala Val Glu Asp Leu Val Ala Ala Lys Leu
219      290                 295                 300
222 Val Tyr Asp Ser Trp Ser Ser Gly Lys
223 305                 310
226 <210> SEQ ID NO: 9
227 <211> LENGTH: 313
228 <212> TYPE: PRT
229 <213> ORGANISM: Mus musculus
231 <400> SEQUENCE: 9
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237 Leu Arg Ser Ser Ser Leu Leu Ile Pro Pro Leu Glu Ala Ala Leu Ala
238             20                  25                  30
241 Asn Phe Ser Lys Gly Pro Asp Gly Gly Val Met Gln Pro Val Arg Thr
242          35                  40                  45
245 Val Val Pro Val Ala Lys His Arg Gly Phe Leu Gly Val Met Pro Ala
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246     50                  55                  60
249 Tyr Ser Ala Ala Glu Asp Ala Leu Thr Thr Lys Leu Val Thr Phe Tyr
250 65                  70                  75                  80
253 Glu Gly His Ser Asn Thr Ala Val Pro Ser His Gln Ala Ser Val Leu
254                 85                  90                  95
257 Leu Phe Asp Pro Ser Asn Gly Ser Leu Leu Ala Val Met Asp Gly Asn
258             100                 105                 110
261 Val Ile Thr Ala Lys Arg Thr Ala Ala Val Ser Ala Ile Ala Thr Lys
262         115                 120                 125
265 Leu Leu Lys Pro Pro Gly Ser Asp Val Leu Cys Ile Leu Gly Ala Gly
266     130                 135                 140
269 Val Gln Ala Tyr Ser His Tyr Glu Ile Phe Thr Glu Gln Phe Ser Phe
270 145                 150                 155                 160
273 Lys Glu Val Arg Met Trp Asn Arg Thr Arg Glu Asn Ala Glu Lys Phe
274                 165                 170                 175
277 Ala Ser Thr Val Gln Gly Asp Val Arg Val Cys Ser Ser Val Gln Glu
278             180                 185                 190
281 Ala Val Thr Gly Ala Asp Val Ile Ile Thr Val Thr Met Ala Thr Glu
282         195                 200                 205
285 Pro Ile Leu Phe Gly Glu Trp Val Lys Pro Gly Ala His Ile Asn Ala
286     210                 215                 220
289 Val Gly Ala Ser Arg Pro Asp Trp Arg Glu Leu Asp Asp Glu Leu Met
290 225                 230                 235                 240
293 Arg Gln Ala Val Leu Tyr Val Asp Ser Arg Glu Ala Ala Leu Lys Glu
294                 245                 250                 255
297 Ser Gly Asp Val Leu Leu Ser Gly Ala Asp Ile Phe Ala Glu Leu Gly
298             260                 265                 270
301 Glu Val Ile Ser Gly Ala Lys Pro Ala His Cys Glu Lys Thr Thr Val
302         275                 280                 285
305 Phe Lys Ser Leu Gly Met Ala Val Glu Asp Leu Val Ala Ala Lys Leu
306     290                 295                 300
309 Val Tyr Asp Ser Trp Ser Ser Gly Lys
310 305                 310
313 <210> SEQ ID NO: 10
314 <211> LENGTH: 314
315 <212> TYPE: PRT
316 <213> ORGANISM: Homo sapiens
318 <400> SEQUENCE: 10
320 Met Ser Arg Val Pro Ala Phe Leu Ser Ala Ala Glu Glu Glu Asp His
321 1               5                   10                  15
324 Leu Arg Ser Ser Ser Leu Leu Ile Pro Pro Leu Glu Thr Ala Leu Ala
325             20                  25                  30
328 Asn Phe Ser Ser Gly Glu Asp Gly Gly Val Met Gln Pro Val Arg Thr
329         35                  40                  45
332 Val Val Pro Val Thr Lys His Arg Gly Tyr Leu Gly Val Met Pro Ala
333     50                  55                  60
336 Tyr Ser Ala Ala Glu Asp Ala Leu Thr Thr Lys Leu Val Thr Phe Tyr
337 65                  70                  75                  80
340 Glu Asp Arg Gly Ile Thr Ser Val Val Pro Ser His Gln Ala Thr Val
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341                 85                  90                  95
344 Leu Leu Phe Glu Pro Ser Asn Gly Thr Leu Leu Ala Val Met Asp Gly
345             100                 105                 110
348 Asn Val Ile Thr Ala Lys Arg Thr Ala Ala Val Ser Ala Ile Ala Thr
349         115                 120                 125
352 Lys Phe Leu Lys Pro Pro Ser Ser Glu Val Leu Cys Ile Leu Gly Ala
353     130                 135                 140
356 Gly Val Gln Ala Tyr Ser His Tyr Glu Ile Phe Thr Glu Gln Phe Ser
357 145                 150                 155                 160
360 Phe Lys Glu Val Arg Ile Trp Asn Arg Thr Lys Glu Asn Ala Glu Lys
361                 165                 170                 175
364 Phe Ala Asp Thr Val Gln Gly Glu Val Arg Val Cys Ser Ser Val Gln
365             180                 185                 190
368 Glu Ala Val Ala Gly Ala Asp Val Ile Ile Thr Val Thr Leu Ala Thr
369         195                 200                 205
372 Glu Pro Ile Leu Phe Gly Glu Trp Val Lys Pro Gly Ala His Ile Asn
373     210                 215                 220
376 Ala Val Gly Ala Ser Arg Pro Asp Trp Arg Glu Leu Asp Asp Glu Leu
377 225                 230                 235                 240
380 Met Glu Gln Ala Val Leu Tyr Val Asp Ser Gln Glu Ala Ala Leu Lys
381                 245                 250                 255
384 Glu Ser Gly Asp Val Leu Leu Ser Gly Ala Glu Ile Phe Ala Glu Leu
385             260                 265                 270
388 Gly Glu Val Ile Lys Gly Val Lys Pro Ala His Cys Glu Lys Thr Thr
389         275                 280                 285
392 Val Phe Lys Ser Leu Gly Met Ala Val Glu Asp Thr Val Ala Ala Lys
393     290                 295                 300
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397 305                 310
400 <210> SEQ ID NO: 11
401 <211> LENGTH: 1015
402 <212> TYPE: DNA
403 <213> ORGANISM: Rattus norvegicus
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408 gagatttctg aaagaaaact gccgtccaga tggaagagaa cttggtgaat tcagaaccac    120
410 aactgtcaac ataggttcga tcagtacagc ggatggctct gctctagtga agctggggaa    180
412 caccacagtc atttgtggag ttaaagcaga atttgcagca ccaccagtag atgcccctga    240
414 tagaggatat gtcgtcccta atgtggacct accaccgctg tgttcatcga ggtttcggac    300
416 tggacctcct ggagaagagg ctcaagtaac cagccagttc attgcagatg tcattgagaa    360
418 ctcacacata attaagaaag aggacttatg catttctcca gggaagcttg cttgggttct    420
420 atactgtgac cttatttgcc tagactacga tgggaacatt ttggatgcct gcacatttgc    480
422 tttgttagca gctttaaaga atgtacagtt gcctgaagtt actataaatg aagaaactgc    540
424 tttagcggaa gtcaatttaa agaagaaaag ttatttgaat gttagagcaa acccagttgc    600
426 tacttcattt gctgtgtttg atgacacttt gctgatagtc gatcctaccg gggaggaggg    660
428 gcaccctgtc cacaggaacc ttaaccgtag taatggacga ggaaggcaag ctgtgctgtc    720
430 ttcacaagcc aggtgggagt gggctgctgg agctaaactt caggactgca tgagtcgagc    780
432 agtaacgaga cacaaagaag tgagcaaact actggatgaa gtaattcaga gcatgaaaca    840
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VERIFICATION SUMMARY DATE: 03/15/2005
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